

RAW SEQUENCE LISTING

**The Biotechnology Systems Branch of the Scientific and Technical
Information Center (STIC) no errors detected.**

Application Serial Number: 09/657, 431 A
Source: TFW16
Date Processed by STIC: 08/03/2005

ENTERED



IFW16

RAW SEQUENCE LISTING

DATE: 08/03/2005

PATENT APPLICATION: US/09/657,431A

TIME: 08:18:41

Input Set : A:\SEQLIST500862001400.TXT

Output Set: N:\CRF4\08032005\I657431A.raw

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4 <110> APPLICANT: ConjuChem, Inc.
5     Beliveau, Richard
6     Bridon, Dominique
7     Rasamoelisololo, Michele
8     Thibaudeau, Karen
9     Huang, Xicai
11 <120> TITLE OF INVENTION: Long Lasting Anti-Angiogenic Peptides
14 <130> FILE REFERENCE: 500862001400
16 <140> CURRENT APPLICATION NUMBER: 09/657,431A
17 <141> CURRENT FILING DATE: 2000-09-07
19 <150> PRIOR APPLICATION NUMBER: 60/153,406
20 <151> PRIOR FILING DATE: 1999-09-10
22 <150> PRIOR APPLICATION NUMBER: 60/159,783
23 <151> PRIOR FILING DATE: 1999-10-15
25 <160> NUMBER OF SEQ ID NOS: 41
27 <170> SOFTWARE: FastSEQ for Windows Version 4.0
29 <210> SEQ ID NO: 1
30 <211> LENGTH: 790
31 <212> TYPE: PRT
32 <213> ORGANISM: Artificial Sequence
34 <220> FEATURE:
35 <223> OTHER INFORMATION: Synthetic Peptide
37 <400> SEQUENCE: 1
38 Glu Pro Leu Asp Asp Tyr Val Asn Thr Gln Gly Ala Ser Leu Phe Ser
39 1           5           10           15
40 Val Thr Lys Lys Gln Leu Gly Ala Gly Ser Ile Glu Glu Cys Ala Ala
41           20           25           30
42 Lys Cys Glu Glu Asp Glu Glu Phe Thr Cys Arg Ala Phe Gln Tyr His
43           35           40           45
44 Ser Lys Glu Gln Gln Cys Val Ile Met Ala Glu Asn Arg Lys Ser Ser
45           50           55           60
46 Ile Ile Ile Arg Met Arg Asp Val Val Leu Phe Glu Lys Lys Val Tyr
47 65           70           75           80
48 Leu Ser Glu Cys Lys Thr Gly Asn Gly Lys Asn Tyr Arg Gly Thr Ser
49           85           90           95
50 Lys Thr Lys Asn Gly Ile Thr Cys Gln Lys Trp Ser Ser Thr Ser Pro
51           100          105          110
52 His Arg Pro Arg Phe Ser Pro Ala Thr His Pro Ser Glu Gly Leu Glu
53           115          120          125
54 Glu Asn Tyr Cys Arg Asn Pro Asp Asn Asp Pro Gln Gly Pro Trp Cys
55           130          135          140
56 Tyr Thr Thr Asp Pro Glu Lys Arg Tyr Asp Tyr Cys Asp Ile Leu Glu
57 145          150          155          160

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58 Cys Glu Glu Glu Cys Met His Cys Ser Gly Glu Asn Tyr Asp Gly Lys
59          165          170          175
60 Ile Ser Lys Thr Met Ser Gly Leu Glu Cys Gln Ala Trp Asp Ser Gln
61          180          185          190
62 Ser Pro His Ala His Gly Tyr Ile Pro Ser Lys Phe Pro Asn Lys Asn
63          195          200          205
64 Leu Lys Lys Asn Tyr Cys Arg Asn Pro Asp Arg Glu Leu Arg Pro Trp
65          210          215          220
66 Cys Phe Thr Thr Asp Pro Asn Lys Arg Trp Glu Leu Cys Asp Ile Pro
67 225          230          235          240
68 Arg Cys Thr Thr Pro Pro Pro Ser Ser Gly Pro Thr Tyr Gln Cys Leu
69          245          250          255
70 Lys Gly Thr Gly Glu Asn Tyr Arg Gly Asn Val Ala Val Thr Val Ser
71          260          265          270
72 Gly His Thr Cys Gln His Trp Ser Ala Gln Thr Pro His Thr His Asn
73          275          280          285
74 Arg Thr Pro Glu Asn Phe Pro Cys Lys Asn Leu Asp Glu Asn Tyr Cys
75          290          295          300
76 Arg Asn Pro Asp Gly Lys Arg Ala Pro Trp Cys His Thr Thr Asn Ser
77 305          310          315          320
78 Gln Val Arg Trp Glu Tyr Cys Lys Ile Pro Ser Cys Asp Ser Ser Pro
79          325          330          335
80 Val Ser Thr Glu Gln Leu Ala Pro Thr Ala Pro Pro Glu Leu Thr Pro
81          340          345          350
82 Val Val Gln Asp Cys Tyr His Gly Asp Gly Gln Ser Tyr Arg Gly Thr
83          355          360          365
84 Ser Ser Thr Thr Thr Thr Gly Lys Lys Cys Gln Ser Trp Ser Ser Met
85          370          375          380
86 Thr Pro His Arg His Gln Lys Thr Pro Glu Asn Tyr Pro Asn Ala Gly
87 385          390          395          400
88 Leu Thr Met Asn Tyr Cys Arg Asn Pro Asp Ala Asp Lys Gly Pro Trp
89          405          410          415
90 Cys Phe Thr Thr Asp Pro Ser Val Arg Trp Glu Tyr Cys Asn Leu Lys
91          420          425          430
92 Lys Cys Ser Gly Thr Glu Ala Ser Val Val Ala Pro Pro Pro Val Val
93          435          440          445
94 Leu Leu Pro Asp Val Glu Thr Pro Ser Glu Glu Asp Cys Met Phe Gly
95          450          455          460
96 Asn Gly Lys Gly Tyr Arg Gly Lys Arg Ala Thr Thr Val Thr Gly Thr
97 465          470          475          480
98 Pro Cys Gln Asp Trp Ala Ala Gln Glu Pro His Arg His Ser Ile Phe
99          485          490          495
100 Thr Pro Glu Thr Asn Pro Arg Ala Gly Leu Glu Lys Asn Tyr Cys Arg
101          500          505          510
102 Asn Pro Asp Gly Asp Val Gly Gly Pro Trp Cys Tyr Thr Thr Asn Pro
103          515          520          525
104 Arg Lys Leu Tyr Asp Tyr Cys Asp Val Pro Gln Cys Ala Ala Pro Ser
105          530          535          540
106 Phe Asp Cys Gly Lys Pro Gln Val Glu Pro Lys Lys Cys Pro Gly Arg

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107 545          550          555          560
108 Val Val Gly Gly Cys Val Ala His Pro His Ser Trp Pro Trp Gln Val
109          565          570          575
110 Ser Leu Arg Thr Arg Phe Gly Met His Phe Cys Gly Gly Thr Leu Ile
111          580          585          590
112 Ser Pro Glu Trp Val Leu Thr Ala Ala His Cys Leu Glu Lys Ser Pro
113          595          600          605
114 Arg Pro Ser Ser Tyr Lys Val Ile Leu Gly Ala His Gln Glu Val Asn
115          610          615          620
116 Leu Glu Pro His Val Gln Glu Ile Glu Val Ser Arg Leu Phe Leu Glu
117 625          630          635          640
118 Pro Thr Arg Lys Asp Ile Ala Leu Leu Lys Leu Ser Ser Pro Ala Val
119          645          650          655
120 Ile Thr Asp Lys Val Ile Pro Ala Cys Leu Pro Ser Pro Asn Tyr Val
121          660          665          670
122 Val Ala Asp Arg Thr Glu Cys Phe Ile Thr Gly Trp Gly Glu Thr Gln
123          675          680          685
124 Gly Thr Phe Gly Ala Gly Leu Leu Lys Glu Ala Gln Leu Pro Val Ile
125          690          695          700
126 Glu Asn Lys Val Cys Asn Arg Tyr Glu Phe Leu Asn Gly Arg Val Gln
127 705          710          715          720
128 Ser Thr Glu Leu Cys Ala Gly His Leu Ala Gly Gly Thr Asp Ser Cys
129          725          730          735
130 Gln Gly Asp Ser Gly Gly Pro Leu Val Cys Phe Glu Lys Asp Lys Tyr
131          740          745          750
132 Ile Leu Gln Gly Val Thr Ser Trp Gly Leu Gly Cys Ala Arg Pro Asn
133          755          760          765
134 Lys Pro Gly Val Tyr Val Arg Val Ser Arg Phe Val Thr Trp Ile Glu
135          770          775          780
136 Gly Val Met Arg Asn Asn
137 785          790
140 <210> SEQ ID NO: 2
141 <211> LENGTH: 7
142 <212> TYPE: PRT
143 <213> ORGANISM: Artificial Sequence
145 <220> FEATURE:
146 <223> OTHER INFORMATION: Synthetic Peptide
148 <400> SEQUENCE: 2
149 Pro Arg Lys Leu Tyr Asp Lys
150 1          5
153 <210> SEQ ID NO: 3
154 <211> LENGTH: 12
155 <212> TYPE: PRT
156 <213> ORGANISM: Artificial Sequence
158 <220> FEATURE:
159 <223> OTHER INFORMATION: Synthetic Peptide
161 <400> SEQUENCE: 3
162 Tyr Thr Thr Asn Pro Arg Lys Leu Tyr Asp Tyr Lys
163 1          5          10

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166 <210> SEQ ID NO: 4
167 <211> LENGTH: 24
168 <212> TYPE: PRT
169 <213> ORGANISM: Artificial Sequence
171 <220> FEATURE:
172 <223> OTHER INFORMATION: Synthetic Peptide
174 <400> SEQUENCE: 4
175 Arg Asn Pro Asp Gly Asp Val Gly Gly Pro Trp Ala Tyr Thr Thr Asn
176 1 5 10 15
177 Pro Arg Lys Leu Tyr Asp Tyr Lys
178 20
181 <210> SEQ ID NO: 5
182 <211> LENGTH: 12
183 <212> TYPE: PRT
184 <213> ORGANISM: Artificial Sequence
186 <220> FEATURE:
187 <223> OTHER INFORMATION: Synthetic Peptide
189 <400> SEQUENCE: 5
190 Arg Asn Pro Asp Gly Asp Val Gly Gly Pro Trp Lys
191 1 5 10
194 <210> SEQ ID NO: 6
195 <211> LENGTH: 8
196 <212> TYPE: PRT
197 <213> ORGANISM: Artificial Sequence
199 <220> FEATURE:
200 <223> OTHER INFORMATION: Synthetic Peptide
202 <400> SEQUENCE: 6
203 Pro Arg Lys Leu Tyr Asp Tyr Lys
204 1 5 10
207 <210> SEQ ID NO: 7
208 <211> LENGTH: 12
209 <212> TYPE: PRT
210 <213> ORGANISM: Artificial Sequence
212 <220> FEATURE:
213 <223> OTHER INFORMATION: Synthetic Peptide
215 <400> SEQUENCE: 7
216 Tyr Thr Thr Asn Pro Arg Lys Leu Tyr Asp Tyr Lys
217 1 5 10
220 <210> SEQ ID NO: 8
221 <211> LENGTH: 11
222 <212> TYPE: PRT
223 <213> ORGANISM: Artificial Sequence
225 <220> FEATURE:
226 <223> OTHER INFORMATION: Synthetic Peptide
228 <400> SEQUENCE: 8
229 Tyr Thr Thr Asn Pro Arg Lys Leu Tyr Asp Tyr
230 1 5 10
233 <210> SEQ ID NO: 9
234 <211> LENGTH: 23

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235 <212> TYPE: PRT
236 <213> ORGANISM: Artificial Sequence
238 <220> FEATURE:
239 <223> OTHER INFORMATION: Synthetic Peptide
241 <400> SEQUENCE: 9
242 Arg Asn Pro Asp Gly Asp Val Gly Gly Pro Trp Ala Tyr Thr Thr Asn
243 1          5          10          15
244 Pro Arg Lys Leu Tyr Asp Tyr
245          20
248 <210> SEQ ID NO: 10
249 <211> LENGTH: 7
250 <212> TYPE: PRT
251 <213> ORGANISM: Artificial Sequence
253 <220> FEATURE:
254 <223> OTHER INFORMATION: Synthetic Peptide
256 <400> SEQUENCE: 10
257 Arg Lys Leu Tyr Asp Tyr Lys
258 1          5
261 <210> SEQ ID NO: 11
262 <211> LENGTH: 6
263 <212> TYPE: PRT
264 <213> ORGANISM: Artificial Sequence
266 <220> FEATURE:
267 <223> OTHER INFORMATION: Synthetic Peptide
269 <400> SEQUENCE: 11
270 Arg Lys Leu Tyr Asp Tyr
271 1          5
274 <210> SEQ ID NO: 12
275 <211> LENGTH: 7
276 <212> TYPE: PRT
277 <213> ORGANISM: Artificial Sequence
279 <220> FEATURE:
280 <223> OTHER INFORMATION: Synthetic Peptide
282 <400> SEQUENCE: 12
283 Pro Arg Lys Leu Tyr Asp Lys
284 1          5
287 <210> SEQ ID NO: 13
288 <211> LENGTH: 6
289 <212> TYPE: PRT
290 <213> ORGANISM: Artificial Sequence
292 <220> FEATURE:
293 <223> OTHER INFORMATION: Synthetic Peptide
295 <400> SEQUENCE: 13
296 Pro Arg Lys Leu Tyr Asp
297 1          5
300 <210> SEQ ID NO: 14
301 <211> LENGTH: 8
302 <212> TYPE: PRT
303 <213> ORGANISM: Artificial Sequence

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RAW SEQUENCE LISTING ERROR SUMMARY
PATENT APPLICATION: US/09/657,431A

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Input Set : A:\SEQLIST500862001400.TXT
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Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:17; Xaa Pos. 1,7
Seq#:18; Xaa Pos. 1,7
Seq#:19; Xaa Pos. 1,12
Seq#:20; Xaa Pos. 1,24
Seq#:21; Xaa Pos. 1,12
Seq#:22; Xaa Pos. 1,8
Seq#:23; Xaa Pos. 1,7
Seq#:24; Xaa Pos. 1,7
Seq#:25; Xaa Pos. 1,12
Seq#:26; Xaa Pos. 1,11
Seq#:27; Xaa Pos. 1,11
Seq#:28; Xaa Pos. 1,24
Seq#:29; Xaa Pos. 1,23
Seq#:30; Xaa Pos. 1,23
Seq#:31; Xaa Pos. 1,12
Seq#:32; Xaa Pos. 1,11
Seq#:33; Xaa Pos. 1,11
Seq#:34; Xaa Pos. 1,7
Seq#:35; Xaa Pos. 1,6
Seq#:36; Xaa Pos. 1,6
Seq#:37; Xaa Pos. 1,7
Seq#:38; Xaa Pos. 1,6
Seq#:39; Xaa Pos. 1,6
Seq#:40; Xaa Pos. 1,8
Seq#:41; Xaa Pos. 1,8

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Input Set : A:\SEQLIST500862001400.TXT

Output Set: N:\CRF4\08032005\I657431A.raw

L:347 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:351 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:17
L:355 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:17
L:356 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:17 after pos.:0
L:368 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:372 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:18
L:376 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:18
L:377 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:18 after pos.:0
L:389 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:393 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:19
L:397 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:19
L:398 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:19 after pos.:0
L:410 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:414 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:20
L:418 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:20
L:419 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:20 after pos.:0
M:341 Repeated in SeqNo=20
L:433 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:437 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:21
L:441 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:21
L:442 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:21 after pos.:0
L:454 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:458 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:22
L:462 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:22
L:463 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:22 after pos.:0
L:475 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:479 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:23
L:483 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:23
L:484 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:23 after pos.:0
L:496 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:500 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:24
L:504 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:24
L:505 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:24 after pos.:0
L:517 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:521 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:25
L:525 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:25
L:526 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:25 after pos.:0
L:538 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:542 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:26
L:546 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:26
L:547 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:26 after pos.:0
L:559 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:563 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:27
L:567 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:27
L:568 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27 after pos.:0
L:580 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:584 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:28
L:588 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:28

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L:589 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:28 after pos.:0
M:341 Repeated in SeqNo=28
L:603 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:607 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:29
L:611 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:29
L:612 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:29 after pos.:0
M:341 Repeated in SeqNo=29
L:626 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:630 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:30
L:634 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:30
L:635 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:30 after pos.:0
M:341 Repeated in SeqNo=30
L:649 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:653 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:31
L:657 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:31
L:658 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:31 after pos.:0
L:670 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:674 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:32
L:678 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:32
L:679 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:32 after pos.:0
L:691 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:695 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:33
L:699 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:33
L:700 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:33 after pos.:0
L:712 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:716 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:34
L:720 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:34
L:721 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:34 after pos.:0
L:733 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:737 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:35
L:741 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:35
L:742 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:35 after pos.:0
L:754 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:758 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:36
L:762 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:36
L:763 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:36 after pos.:0
L:775 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:779 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:37
L:783 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:37
L:784 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:37 after pos.:0
L:796 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:800 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:38
L:804 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:38
L:805 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:38 after pos.:0
L:817 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:821 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:39
L:825 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:39
L:826 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:39 after pos.:0
L:838 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!

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L:842 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:40
L:846 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:40
L:847 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:40 after pos.:0
L:859 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:863 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:41
L:868 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:41
L:869 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:41 after pos.:0